

**Controlling Offspring's Sex Ratio by Targeting
Transgenes onto the Sex Chromosomes**

Cross-Reference to Related Application

- 5 This application claims the priority benefit of U.S. provisional application serial no. 60/173,096, filed December 27, 1999, of the same title and by the same inventors.

Background of the Invention

- 10 1. Field of the Invention

The present invention relates to methods and compositions of matter useful for making transgenic animals whose offspring's sex ratio can be altered through the expression of said transgene integrated onto one of the two sex chromosomes.

- 15 2. Description of the Prior Art

The majority of animal species consists of two sexes, male and female. For many species, males and females are different in many aspects such as body size, growth rate and behavior. Certain features are even unique to one sex. In farm animals, one sex may be advantageous over the other for producing desired products such as meat, milk, egg, and wool. Thus, controlling the birth ratio of the two sexes of livestock is economically important, since it allows farmers to take full advantage of the differences between the two sexes.

In mammals, as well as in many other organisms, sex is determined by the sex chromosome (X or Y) of the sperm. Normally, an egg fertilized by a sperm that contains the Y chromosome (Y-sperm) will develop into a male, and on the other hand, an egg fertilized by a sperm that contains the X chromosome (X-sperm) will become a female. This genetic mechanism determines that approximately equal numbers of males and females are born. The key for controlling sex is to select the right sperm (X or Y) for fertilization. During the past three decades, several immunological, mechanical, and chemical methods have been tested to separate the X and Y sperm from collected semen. The idea is to use separated sperm to fertilize eggs by artificial insemination or in vitro fertilization with the hope to change the sex

ratio of the offspring. Although dozens of patents have been issued to cover these techniques, these methods are generally considered inefficient and unreliable. Further, they require equipment and technical expertise that are not available to most farmers. Recently, Sry (Sex-determine Region Y) gene has been identified as
5 an important male determinant in mammals. Several laboratories have tried to preferentially increase the percentage of male newborns by expressing Sry gene in transgenic animals. However, only 30% of the XX transgenic mice (normally female) grew male organs such as testis, and these sex-reversed animals were sterile.

10 Therefore, what is needed is a novel approach to entirely eliminate the production of either X or Y sperm in the testis so that all the offspring of such male will be the same sex, either male or female. What is also needed is such an approach, such that the animals are genetically modified so that this special trait can be passed from one generation to another. Further, what is needed is such an
15 approach that is distinct from trying to separate the X and Y sperm in the semen. Such an approach should control sex ratio more precisely, and also eliminate the tricky separation process for each semen sample.

Summary of the Invention

20 Here, we developed a novel approach that could entirely eliminate the production of either X or Y sperm in the testis. Therefore, all the offspring of such male will be the same sex, either male or female. The animals are genetically modified so that this special trait can be passed from one generation to another. This approach is clearly better than those trying to separate the X and Y sperm in the
25 semen. It can control sex ratio more precisely, and also eliminate the tricky separation process for each semen sample.

The method of the present invention involves: (a) selecting or creating a transgene whose expression can interfere with sperm's ability to undergo
30 fertilization, and whose gene products (mRNA and protein) do not diffuse freely among inter-connected spermatids; (b) placing the said transgene under the regulatory control of post-meiotic spermatogenesis-specific promoter; (c) using the said transgene to generate transgenic animals in the way that the transgene is

inserted onto one of the two sex chromosomes. Expression of the said transgene can reduce the ability of the sperm with one particular sex chromosome to fertilize eggs and develop into new individuals, and consequently controlling the sex ratio of the offspring. This method can also be modified by replacing the post-meiotic 5 spermatogenesis-specific promoter with a promoter that expresses during embryogenesis (after fertilization), and replacing the transgene that is toxic to sperm with a transgene that is toxic to early embryos. Embryonic expression of a sex chromosome-linked toxin transgene disrupts the normal development of embryos with one particular sex chromosome, and only allows embryos with preferred sex to 10 develop into viable individuals. This modified method can be used not only in organisms using X and Y chromosomes but also Z and W chromosomes for sex determination. This method allows farmers to choose the preferred sex of farm animals to produce dairy and meat products and allows laboratory animal suppliers to specifically increase the birth ratio of the preferred gender.

15 These and other advantages of the present invention will become apparent upon review of the following description, the accompanying drawings, and the appended claims.

Brief Description of the Drawing

20 The FIGURE is a simplified representation of a method of the present invention for altering the sex chromosomes in animals.

Detailed Description of the Invention

25 Methods

1. Rationale and strategy

It has been demonstrated in transgenic mice that a tissue or a cell type can 30 be specifically eliminated by expressing a toxic gene product. Therefore, post-meiotic expression of a toxin gene in the haploid spermatids destroys the sperm that contain the toxin transgene. When the transgene is integrated into one of the two sex chromosomes (X or Y), only the spermatids containing that particular sex

chromosome are disabled while the spermatids containing the other sex chromosome may function normally. Such male transgenic animals can only produce one type of sperm, and hence all of their offspring should be the same sex.

A major challenge to the above theory is the fact that the haploid spermatids remain

5 connected by cytoplasmic bridges (~1 um), i.e. they are syncytial. Therefore, the toxin could diffuse into the spermatids that do not contain the transgene, leading to the destruction of all sperm rather than only the half with the toxin transgene (Braun et. al., 1989, see review Davies and Willison, 1993). However, it is possible to solve this problem by choosing a toxin gene whose products can not diffuse freely among

10 the inter-connected spermatids, or by creating transgenes with limited diffusion characteristics by combining cellular localization DNA sequences to the toxin transgenes. A technique for achieving that is represented in a simplified manner in the accompanying FIGURE.

15 In the current invention, a herpes simplex type-1 virus thymidine kinase (HSV-tk) gene is used as the toxin transgenes. It has been previously shown that HSV-tk contains a spermatogenesis-specific cryptic promoter and its expression disrupt spermatogenesis (Palmiter et al. 1984; Ellison and Bishop, 1988; Braun et al., 1990; Wilkie et al., 1991; Al-Shawi et al., 1991; Huttner et al., 1993; Bernier et al., 1994; Salomon et al., 1995; Ellison et al., 1995; Ellison and Bishop, 1996; Al-Shawi et al., 1998). One common phenotype for all these HSV-tk transgenic mice is that the male mice can not transmit the HSV-tk transgene to the next generation although many of these males are fertile. Gondo et al. (1994) It has been reported that embryonic stem (ES) cell lines containing the HSV-tk transgene could not transmit

20 through the male germ-line either. All these results together with the report that HSV-tk protein was preferentially distributed at the perinuclear region of transfected cell (Haarr and Flatmark, 1987) implies that the diffusion of HSV-tk gene products (mRNA and protein) among the inter-connected spermatids is limited. Therefore, HSV-tk is an ideal candidate for use in the sex control project. Of course, it is

25 possible to find other genes with similar properties. Alternatively, such toxin genes can be created by linking cellular localization DNA sequences to a variety of genes that may interfere with sperm's function, such as genes that cause cell death, regulatory genes that block sperm development and maturation, mutated

cytoskeleton or energy-producing genes involved in sperm motility, and genes involved in gamete recognition, penetration and fusion.

The toxic transgene can be inserted onto the sex chromosome by a variety of methods, and several examples are briefly described below.

(1) Gene-targeting using embryonic stem (ES) cells technology

(a) Targeting the transgene onto the sex chromosome of ES cells using the standard homologous recombination method.

(b) Microinjecting the positive ES cells into recipient embryos to make chimeric mice.

(c) Breeding the chimeric mice to pass the ES-derived germ cells to the next generation to obtain the desired transgenic animals.

(2) Gene-targeting using animal cloning technology

(a) Targeting the transgene onto the sex chromosome of suitable nuclear donor cells (such as fibroblasts derived from embryos) by using the standard homologous recombination method.

(b) Transferring the nuclei of the donor cells into enucleated oocytes by cell fusion or nuclear transfer.

(c) Activating the reconstituted oocytes and transferring them into pseudo pregnant foster mothers to allow the embryos to develop into individuals with the transgene inserted onto the desired chromosome.

(3) Random integration using ES cell or cloning technology

(a) Co-transfecting the HSV-tk transgene with a selection marker gene (such as neomycin-resistant gene) into ES or nuclear donor cells by various transfection methods (such as electroporation).

(b) Picking and growing neomycin-resistant clones, and examining the transgene integration site by fluorescence in situ hybridization (FISH). Expanding the clones with the transgene inserted onto the desired sex chromosome.

(c) Injecting the ES cells into embryos to make chimeric animals, or cloning the animals using the donor cells' nuclei.

(4) Random insertion of the transgene into gametes or germ cells

Using a variety of methods such as pronuclear injection, retroviral vector transfection, lipofection, and sperm incubation to randomly insert the transgene into the genome. Then, using FISH to screen the transgenic founders to identify individuals with the transgene inserted onto the desired sex chromosome.

5

Methods (3) and (4) use a random integration mechanism. The advantage for these methods is they are relatively easy to use to make the DNA constructs. The disadvantage is it is necessary to screen a considerable number of cell clones or individuals in order to find the one with the transgene inserted onto the desired sex 10 chromosome. Methods (1) and (2) use a homologous recombination method to specifically target the transgene onto the desired sex chromosome, and therefore eliminate the need for screening using FISH. However, it involves more to make the gene-targeting DNA constructs. In addition to the selectable marker genes that need to be incorporated into the targeting construct, two flanking homologous DNA 15 fragments are also necessary for the homologous recombination events to occur. As more and more DNA sequence data are available from the sex chromosomes, many loci can be used as target sites for inserting the transgene. The general rules are that the target loci should be accessible by the transcription machinery for transcription in spermatids and the insertion of the transgene into those sites do not cause abnormal phenotype to the transgenic animals.

Since the transgenic males can not transmit the HSV-tk transgene from one generation to the next, an inducible (such as the tetracyclin inducible system) or conditional system (such as Cre/loxP) need to be used. For example, loxP sites 25 flanked by intervening DNA sequence can be inserted between the promoter and the transgene to prevent the expression of the HSV-tk transgene. Therefore, the transgene can be transmitted through generations to maintain the lines. When needed for sex control, the HSV-tk transgene can be activated by removing the intervening DNA sequences (between the two loxP sites) through mating with 30 females containing the Cre recombinase gene. The Cre transgenic animals can be purchased through commercial sources or can be created using standard transgenic methods. For transgenes targeted on the X chromosome, it is possible to maintain the transgenic line through females (mother to daughter). Therefore, it is not

necessary to use the inducible or conditional systems after the line is created.

2. Modified method

The above technology can be modified by replacing the post-meiotic spermatogenesis-specific promoters with promoters that express during embryogenesis (after fertilization), and replacing the toxin transgene that disrupts the sperm's function to a transgene that affects embryonic development or viability. When such transgenes are inserted onto one of the two sex chromosomes, the expression of the transgenes during early embryogenesis disrupts the normal development of embryos with one particular sex chromosome, and allows only embryos with preferred sex to develop into viable individuals. This modified method will reduce the litter size by a half. However, if it is combined with superovulation (injecting hormones to increase the number of eggs per ovulation), it is a useful method for sex selection.

This modified method needs an inducible or conditional system for maintaining the transgene through generations since all embryos with the transgene are killed and therefore can not transmit to the next generation. For example, the toxin transgene can be prevented from transcription by inserting a loxP site-flanked intervening sequence between the promoter and the transcription unit. Such inactive transgene can be passed from generation to generation. When needed for sex control, the transgenic animals can mate with animals containing a Cre recombinase gene controlled by gametogenesis-specific promoters. The Cre recombinase will activate the transgenes in the gametes, the embryos developed from such gametes will be eliminated, and only the embryos with the sex chromosome that does not contain the transgene can develop into individuals.

The modified method can be used not only in organisms using X and Y but also in organisms using Z and W chromosomes to determine sex. In organisms using a ZW system to determine sex, the females are heterogametic (Z eggs and W eggs) while the males are homogametic (all sperm contain Z chromosome). Therefore, transgenic females are used for sex control. For species which lay a large number of eggs during each ovulation (such as many aquatic species,

amphibians, insects and plants), elimination of half of the early embryos should not significantly influence its reproduction since a majority of the embryos can not develop into adults anyway. However, for those species whose eggs themselves are valuable commodities (such as birds), it needs to be carefully evaluated if the benefit 5 of being able to control the sex of chicks is more than the value of the half of the eggs being destroyed.

3. Practical procedures for proving the method in laboratory mouse

This technology can be used in all organisms that use sex chromosomes for 10 sex determination, but it only needs to be tested in one species to prove the principles. Currently, mouse is the species of choice for proving the method. Below is an example of the specific protocols that can be used to generate the desired transgenic mice.

- 15 1). Purchasing the HSV-tk gene, neomycin-resistant gene (neo), diphtheria toxin (DT) from commercial sources, such as Stratagene.
- 20 2) Making the loxP site DNA by annealing two single-strand oligodeoxynucleotides.
- 25 3) Cloning of the flanking homologous DNA fragments needed for homologous recombination by polymerase chain reaction (PCR) using 129 mouse genomic DNA as a template. For examples, the transgene can be targeted onto the Hprt locus (see Melton et al., 1984 for gene sequences) of the X chromosome, or the Tspy pseudogene locus (see Vogel et al., 1998 for sequences) of the Y chromosome. Specifically, a ~4 Kb fragment can be amplified between exon 6 and exon 7 of the Hprt gene using the primer CAGTACAGCCCCAAATGGT and primer GAGGTCCCTTTCACCAAGCAA; a ~1.3 Kb fragment can be amplified between exon 8 and exon 9 of the Hprt gene using primer AGTTTGTTGGATATGCC and primer CCTCTTAGATGCTGTTACTG; a ~1 Kb DNA fragment can be amplified from the 5' end of the Tspy pseudogene by using primers AGGAGAGTGTGGGCATG and AAATGCACAATCTAAAGC; and a ~1.5 Kb fragment can be amplified from the 3' end of the Tspy pseudogene by using primers CTCCAAGGACTGCTCTCA and AACATGAGAACATGGTA.
- 30 4) Assembling the above DNA fragments according to Figure 1 to make the gene-targeting constructs for X and Y chromosomes.

- 5) Electroporating the DNA constructs into ES cells, and plating the ES cells in medium containing G418 to allow neomycin-resistant colonies to appear. Since the Hprt gene itself is a selectable marker, adding 6-thioguanine (6-TG) in addition to G418 into the selection medium will increase the chance of targeted clones
5 versus randomly integrated clones for the X chromosome project. For the Y chromosome project, the diphtheria toxin (DT) gene in the targeting construct serves as a negative selectable marker, which can also increase the chance of targeted-clones versus randomly integrated-clones.
- 6) Growing the ES cells for 7-10 days in the selection media, and then picking the
10 resistant colonies and expanding them in multi-well culture plates.
- 7) Duplicating the multi-well plates. Cryopreserving one set of plates to store the ES clones, and preparing genomic DNA from the other set of plates.
- 8) Analyzing the genomic DNA samples by PCR and Southern blotting to identify homologous recombination events. ES clones with the transgene targeted onto
15 the correct chromosome locus are further propagated and stored as positive clones.
- 9) Injecting the positive clones into blastocyst stage mouse embryos. Transferring the injected embryos into the uterus of pseudopregnant foster mothers to allow chimeric mice to be born.
- 20 10) Breeding the chimeric mice with wild-type mice to allow transmission of the transgene to the next generation. For the X chromosome project, it is important to breed female chimeras with wild type males for germ-line transmission of the transgene since the transgene can not be transmitted through sperm. For the Y chromosome project, it is necessary to breed male chimeras with wild type females to transmit the inactive form of the transgene.
- 25 11) Screening the offspring of the above mating by PCR or Southern blotting using tail DNA to identify mice with the transgene.
- 12) For the X chromosome project, the transgene can be transmitted through generations by females, and the males are used for sex control since all of their
30 offspring are the same sex.
- 13) For the Y chromosome project, the inactive form of the transgene is maintained through males. When needed for sex control, the transgene can be activated through mating with females containing Cre recombinase activity. The Cre mice

can be purchased from commercial laboratory animal suppliers, or can be made by standard transgenic methods.

The following references provide background information related to the
5 present invention. To the extent any reference is specifically cited in the Detailed Description, it is incorporated herein by reference for the purpose relied upon in describing the invention.

Al-Shawi, R., J. Burke, H. Wallace, C. Jones, S. Harrison, D. Buxton, S. Maley, A. Chandley, and J. O. Bishop (1991) Mol. Cell Biol. 11:4207-4216.

10

Al-Shawi, R., J. Burke, C. T. Jones, J. P. Simons, J. O. Bishop (1988) Mol. Cell. Biol. 8:4821-4828.

15

Bernier F., S. L. Guerin, M. Ouellet, G. Pelletier, and F. Pothier (1994) Transgenics, 1:225-240.

Braun, R. E., R. R. Behringer, J. J. Peschon, R. L. Brinster, and R. D. Palmiter (1989) Nature, 337:373-376.

20

Braun, R. E., D. Lo, C. A. Pinkert, G. Widera, R. A. Flaveill, R. D. Palmiter, and R. L. Brinster (1990) Biol. Reprod. 43:684-693.

Capel, B., A. Swain, S. Nicolis, A. Hacker, M. Walter, P. Koopman, P. Goodfellow, and R. Lovell-Badge (1993) Cell, 73:1019-1030.

25

Davies, P.O. and K. R. Willison (1993) Sem. Dev. Biol., 3:179-188.

Ellison A. R. and J. O. Bishop (1998) Biochim Biophys Acta. 1442:28-38.

30

Ellison, A. R., H. Wallace, R. Al-Shawi, and J. O. Bishop (1995) Mol. Reprod. Dev., 41:425-434.

Ellison, A. R. and J. O. Bishop (1996) Nucleic Acids Res. 24:2073-2079.

Erickson R. P. (1990) *Trens Genet.*, 6:264-269

Flatmark T. and L. Haarr (1987) *J. Gen. Virol.*, 68:2817-2829.

5

Gondo, Y., K. Nakamura, K. Nakao, T. Sasaoka, K. Ito, M. Kimura, and M. Katsuki (1994) *Biochem. Biophys. Res. Commun.* 202:830-837.

Hendriksen, P. J. M., J. W. Hoogerbrugge, A. P. N. Themmen, M. H. M. Koken, J. H. 10 J. Hoeijmakers, B. A. Oostra, T. V. D. Lende, and J. A. grootegoed (1995) *Dev. Biol.* 170:730-733.

Huttner, K. M., J. Pudney, D. S. Milstone, D. Ladd, and J. G. Seidman (1993) *Biol. Reprod.* 49:251-261.

15

Melton, D.W., D. S. Konecki, J. Brennand, and C.T. Caskey (1984) *Proc. Natl. Acad. Sci. USA*, 81:2147-2151.

20

Nagamine, C. M., K. Chan, L. E. Hake, and Y-F. C. Lao (1990) *Genes Dev.*, 4:63-74.
Palmiter, R.D., T. M. Wilkie, H. Y. Chen, and R. L. Brinster (1984) *Cell*, 36: 869-877.

25 Salomon B., S. Maury, L. Loubiere, M. Caruso, R. Onclercq, and D. Klatzman (1995) *Mol. Cell Biol.*, 15:5322-5328.

Shannon, M. and M. A. Handel (1993) *Biol. Reprod.*, 49:770-778.

Vogel, T., H. Boettger-Tong, I. Nanda, F. Dechend, A.I. Agulnik, C.E. Bishop, M. Schmid, and J. Schmidtke (1998) *Chromosome Res.* 6:35-40.

30

Wilkie, T. M., R. E. Braun, W. J. Ehrman, R. D. Palmiter, and R. E. Hammer (1991) *Genes Dev.*, 5:38-48.

The principles and features of the present invention, described in examples above, will be understood more broadly from the following claims. The claims are intended to cover the invention as described and all equivalents.